

This Page Is Inserted by IFW Operations  
and is not a part of the Official Record

## **BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

**IMAGES ARE BEST AVAILABLE COPY.**

**As rescanning documents *will not* correct images,  
please do not report the images to the  
Image Problems Mailbox.**

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization  
International Bureau



(43) International Publication Date  
8 August 2002 (08.08.2002)

PCT

(10) International Publication Number  
**WO 02/060947 A3**

(51) International Patent Classification<sup>7</sup>: **C07K 14/705**,  
A61K 38/17

GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent  
(BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR,  
NE, SN, TD, TG).

(21) International Application Number: PCT/US02/00509

(22) International Filing Date: 18 January 2002 (18.01.2002)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:  
60/265,690 1 February 2001 (01.02.2001) US

(71) Applicant (for all designated States except US): **ELI  
LILLY AND COMPANY** [US/US]; Lilly Corporate  
Center, Indianapolis, IN 46285 (US).

(72) Inventors; and

(75) Inventors/Applicants (for US only): **JENKINS, Nigel**  
[GB/US]; 241 Wyndottee Drive, Carmel, IN 46032 (US).  
**WITCHER, Derrick, Ryan** [US/US]; 10898 Parrot  
Court, Fishers, IN 46038 (US). **WROBLEWSKI, Victor,**  
**John** [US/US]; 1466 Woodpond South Roundabout,  
Carmel, IN 46033 (US).

(74) Agents: **WEBSTER, Thomas, D.** et al.; Eli Lilly and  
Company, Lilly Corporate Center, Indianapolis, IN 46285  
(US).

(81) Designated States (national): AE, AG, AL, AM, AT (util-  
ity model), AU, AZ, BA, BB, BG, BR, BY, BZ, CA,  
CH, CN, CO, CR, CU, CZ (utility model), CZ, DE (util-  
ity model), DE, DK (utility model), DK, DM, DZ, EC, EE  
(utility model), EE, ES, FI (utility model), FI, GB, GD, GE,  
GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ,  
LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN,  
MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD,  
SE, SG, SI, SK (utility model), SK, SL, TJ, TM, TN, TR,  
TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW.

(84) Designated States (regional): ARIPO patent (GH, GM,  
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),  
Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),  
European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR,

**Declarations under Rule 4.17:**

- as to applicant's entitlement to apply for and be granted  
a patent (Rule 4.17(ii)) for the following designations AE,  
AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA,  
CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES,  
FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE,  
KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG,  
MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU,  
SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG,  
UZ, VN, YU, ZA, ZM, ZW, ARIPO patent (GH, GM, KE, LS,  
MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent  
(AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent  
(AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU,  
MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI,  
CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG)
- as to the applicant's entitlement to claim the priority of the  
earlier application (Rule 4.17(iii)) for the following design-  
ations AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY,  
BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC,  
EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN,  
IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV,  
MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,  
PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR,  
TT, TZ, UA, UG, UZ, VN, YU, ZA, ZM, ZW, ARIPO patent  
(GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),  
Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),  
European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB,  
GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ,  
CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD,  
TG)

**Published:**

- with international search report
- before the expiration of the time limit for amending the  
claims and to be republished in the event of receipt of  
amendments

(88) Date of publication of the international search report:  
31 October 2002

For two-letter codes and other abbreviations, refer to the "Guid-  
ance Notes on Codes and Abbreviations" appearing at the begin-  
ning of each regular issue of the PCT Gazette.

(54) Title: GLYCOFORMS A FAS LIGAND INHIBITORY PROTEIN ANALOG

(57) Abstract: The present invention provides FLINT analog isoform compositions having specified average sialic acid content per molecule of FLINT analog.

WO 02/060947 A3

## FLINT ANALOG GLYCOFORMS

## BACKGROUND OF THE INVENTION

FLINT is a glycoprotein involved in regulating  
5 apoptosis. A number of tumor necrosis factor receptor  
proteins ("TNFR proteins") have been isolated in recent  
years, having many potent biological effects. Loss of normal  
activity of these proteins has been implicated in a number  
of disease states.

10 Increased activation of the Fas-FasL signal  
transduction pathway is implicated in a number of  
pathological conditions, including runaway apoptosis (Kondo  
et al., *Nature Medicine* 3(4):409-413 (1997); Galle et al.,  
*J. Exp. Med.* 182:1223-1230 (1995)), and inflammatory disease  
15 resulting from neutrophil activation (Miwa et al., *Nature  
Medicine* 4:1287 (1998)). "Runaway apoptosis" is a level of  
apoptosis greater than normal, or apoptosis occurring at an  
inappropriate time. Pathological conditions caused by  
runaway apoptosis include, for example, organ failure in the  
20 liver, kidneys and pancreas. Inflammatory diseases  
associated with excessive neutrophil activation include  
sepsis, ARDS, SIRS and MODS.

One particular TNFR homologue, referred to herein as  
"FAS Ligand Inhibitory Protein," or "FLINT", binds Fas  
25 Ligand (FasL) thereby preventing the interaction of FasL  
with Fas. FLINT also binds the ligand known as LIGHT to  
prevent the interaction of LIGHT with receptor LTBR, an  
otherwise initiating step in a second, independent apoptotic  
pathway.

30 Compounds such as FLINT can be used to treat or prevent  
diseases or conditions in mammals, including humans, that

clinically may correlate with either, or both, of the binding interactions of Fas to FasL and LIGHT to LTBR.

Many eucaryotic secretory proteins including FLINT are modified with one or more oligosaccharide groups (See PCT applications WO0058466, WO0058465, and WO9950413). The state of glycosylation of such proteins can dramatically affect their physical properties and also be important to protein stability, secretion, and subcellular localization. Moreover, proper glycosylation can be essential for biological activity. In fact, some genes from eucaryotic organisms, when expressed in bacteria (e.g., *E. coli*) yield proteins that have little or no activity by virtue of their lack of glycosylation.

Glycosylation occurs at specific locations along the polypeptide backbone and is usually of two types: O-linked oligosaccharides are attached to serine or threonine residues, while N-linked oligosaccharides are attached to asparagine residues when part of the consensus sequence Asn-X-Ser/Thr, where X can be any amino acid except proline.

Additionally, the structure and composition of N-linked and O-linked oligosaccharides differ. One type of sugar that is commonly found on N-linked and O-linked oligosaccharides is N-acetylneuraminic acid (hereafter referred to as sialic acid). Sialic acid is usually the terminal residue on both N-linked and O-linked oligosaccharides and, by virtue of its negative charge, may confer acidic properties on a glycoprotein.

FLINT has an N-linked glycosylation site at Asn 173 of SEQ ID NO:3 (Asn 144 of SEQ ID NO:1), and O-linked glycosylation sites at Thr 203 (alternatively, Thr 174 of SEQ ID NO:1); and Thr 245 of SEQ ID NO:3 (alternatively, Thr 216 of SEQ ID NO:1). Under ordinary conditions of

recombinant production, the O-linked site at Thr 216 is substantially less glycosylated than the O-linked site at Thr 174 of SEQ ID NO:1.

Sialic acid content has a direct effect on the PK  
5 profile of FLINT analogs. The higher the sialic acid  
content, the slower the clearance from serum *in vivo*. This  
observation has heretofore not been disclosed. The present  
invention relates to the relationship between sialic acid  
content and *in vivo* clearance of FLINT analogs, and to FLINT  
10 analog compositions that have defined level of sialylation.  
Owing to this relationship, under-sialylated FLINT analog is  
cleared more rapidly from the serum of primates than FLINT  
analog with greater sialylation content. Low-sialylated  
FLINT analog is probably cleared from the circulation upon  
15 interaction with certain hepatic receptors, for example, the  
hepatic asialoglycoprotein binding protein (cf. Morrell et  
al. J. Biol. Chem. 243, 155 (1968); Briggs, et al. Am. J.  
Physiol. 227, 1385 (1974); Ashwell et al. Methods Enzymol.  
50, 287 (1978)). Therefore, achieving enhanced levels of  
20 sialylation is expected to improve the therapeutic utility  
of FLINT analogs.

It is an object of the present invention to provide  
isoforms of FLINT analogs having enhanced sialic acid  
content. The isoforms of the present invention are analogs  
25 of FLINT having additional glycosylation sites engineered  
into the native FLINT sequence. These additional  
glycosylation sites provide additional sites for  
sialylation. Pharmaceutical compositions comprising such  
molecules provide FLINT analog compositions with slower  
30 clearance time *in vivo* and enhanced therapeutic benefit.

## SUMMARY OF THE INVENTION

The subject invention relates to FLINT analog sialic acid isoforms. Also provided are methods of preparing FLINT analog isoforms and pharmaceutically acceptable compositions comprising same. This invention also relates to therapeutic methods comprising administering a therapeutically effective amount of these FLINT analog compositions to treat and/or prevent diseases or conditions in mammals including humans.

10       The subject invention relates further to methods of preparing FLINT analog isoforms comprising subjecting material containing FLINT analog to ion exchange chromatography, liquid chromatography, or chromatofocusing, as well as methods to enhance sialylation of FLINT analogs  
15       comprising use of enzymatic processes *in vitro*.

## DETAILED DESCRIPTION OF THE INVENTION

SEQ ID NO:1 - Mature human FLINT, i.e. native FLINT minus the leader sequence.

20       SEQ ID NO:2 - Nucleic acid/cDNA encoding mature human FLINT.

SEQ ID NO:3 - Native human FLINT.

SEQ ID NO:4 - Nucleic acid encoding human FLINT.

25       SEQ ID NO:5 - Oligonucleotide sequence of primer CF119 used in production of analog A12N.

SEQ ID NO:6 - Oligonucleotide sequence of primer CF120 used in production of analog A12N.

SEQ ID NO:7 - Oligonucleotide sequence of primer CF121 used in production of analog A12N.

30       SEQ ID NO:8 - Oligonucleotide sequence of primer CF122 used in production of analog A12N.

The term "analog" or "FLINT analog" is used herein specifically to mean a FLINT variant having one or more amino acid sequence changes in SEQ ID NO:1 or SEQ ID NO:3, e.g. substitution, addition, deletion, such that one or more additional glycosylation site(s) is present when compared with native FLINT. Said analogs or variants retain the biological activity of FLINT.

As used herein, "average sialic acid content" refers to a quantitative measure of the sialic acid content of a FLINT analog sample preparation expressed as the mole fraction of sialic acid per mole of analog. The term allows comparison of different lots or preparations of FLINT analog. FLINT analog preparations may comprise multiple FLINT isoforms, for example, 0, 1, 2, 3, 4, 5, 6 or more sialic acids per molecule of FLINT analog.

The term "native FLINT" refers to SEQ ID NO:3.

The term "mature FLINT" refers to SEQ ID NO:1.

Description of FLINT analog isoforms provided herein may be referenced against SEQ ID NO:1 or SEQ ID NO:3. Both the native and mature forms of FLINT are within the scope of the invention.

"FLINT analog isoforms" refers to sialic acid variants. Native FLINT can occur with 0, 1, 2, 3, 4, 5, or 6 sialic acid residues per molecule of FLINT. Each additional N-linked glycosylation site would provide 0, 1, 2, 3, or 4 additional sialic acids per molecule of analog; each additional O-linked site would provide 0, 1, or 2 additional sialic acids per molecule of analog. The degree of sialylation will depend on the host cell and growth conditions used in producing recombinant FLINT analogs. Sialylation can be enhanced *in vitro* using an enzymatic process described later in this disclosure.

The term "N-glycosylated polypeptide" refers to polypeptides having one or more NXS/T motifs in which the nitrogen atom in the side chain amide of the asparagine is covalently bonded to a glycosyl group. "X" refers to any naturally occurring amino acid residue except proline. The "naturally occurring amino acids" are glycine, alanine, valine, leucine, isoleucine, proline, serine, threonine, cysteine, methionine, lysine, arginine, glutamic acid, aspartic acid, glutamine, asparagine, phenylalanine, histidine, tyrosine and tryptophan. N-Glycosylated proteins are optionally O-glycosylated.

The term "O-glycosylated polypeptide" refers to polypeptides having one or more serines and/or threonine in which the oxygen atom in the side chain is covalently bonded to a glycosyl group. O-Glycosylated proteins are optionally N-glycosylated.

The nucleotide and amino acid abbreviations used herein are those accepted in the art and by the United States Patent and Trademark Office, as set forth in 37 C.F.R. 1.822 (b) (2).

The present invention relates to sialic acid isoforms of analogs of FLINT in which one or more amino acids of native FLINT are substituted, deleted, or added to create additional glycosylation site(s). Preferred analogs include those having one or two additional N-linked glycosylation sites. N-linked sites are created by introducing an N-linked consensus sequence into the native FLINT sequence. N-linked glycosylation consensus sites comprise the sequence NXS/T.

Contemplated by the present invention are specific analogs of SEQ ID NO:1 having substitutions to create new N-linked glycosylation sites at different positions, including: Ala2 or Ala12 to Asn (1 additional N-linked



site); Pro25, Pro38, Pro126, or Pro171 to Asn (1 additional N-linked site); Arg35 to Asn (1 additional N-linked site); Ser37 to Asn and Pro38 to any amino acid (1 additional N-linked site); Ser166 to Asn (1 additional N-linked site);  
5 Leu172 to Asn (1 additional N-linked site); Asp194 to Asn (1 additional N-linked site); Thr114 to Asn and Pro115 to any amino acid (1 additional N-linked site). Also contemplated are substitutions including: Ala12 to Asn and Glu13 to Gln (1 additional N-linked site); Arg34 to Asn and Asp36 to Thr  
10 (1 additional N-linked site); Arg35 to Asn, Ser37 to Thr (1 additional N-linked site); Ser132 to Asn, Ser134 to Thr (1 additional N-linked site); Asp194 to Asn, Ser196 to Thr (1 additional N-linked site); Arg35 and Asp194 to Asn (2 additional N-linked sites); Arg34 to Asn, Asp36 to Thr (1  
15 additional N-linked site); Asp194 to Asn, Ser196 to Thr (1 additional N-linked site); Ser132 to Asn (1 additional N-linked site); or Ala12 and Ser132 to Asn and Ser134 to Thr (1 additional N-linked site); Ala12 to Asn, Glu13 to Gln, Asp 194 to Asn, Ser 196 to Thr (2 additional N-linked  
20 sites); Arg34 to Asn, Asp36 to Thr, Asp194to Asn, Ser196 to Thr (2 additional N-linked sites).

Glycosylation sites may be introduced into the native FLINT cDNA sequence most conveniently by *in vitro* mutagenesis techniques, well known to the skilled artisan  
25 (See e.g., T. Maniatis et al. Molecular Cloning: A Laboratory Manual, 2d Ed.(1989), and Ausubel et al., eds., Current Protocols in Molecular Biology, Wiley Interscience, New York (1987-1999). For example, synthetic  
oligonucleotides are designed to incorporate a point  
30 mutation at one end of an amplified fragment. Following first strand PCR, the amplified fragments encompassing the mutation are annealed with each other and extended by

mutually primed synthesis. Annealing is followed by a second PCR step utilizing 5' forward and 3' reverse end primers in which the entire mutagenized fragment gets amplified and is ready for subcloning into an appropriate vector.

- 5        Analogs are generated by site-directed mutagenesis and include additions, deletions, or substitutions of amino acid residues that add sites that are available for glycosylation. Analogs of the invention potentially have a greater number of carbohydrate chains than native FLINT, 10 their sequence having additional N-linked glycosylation sites engineered therein.

- FLINT analogs were constructed using the method described in Nelson, R. M. and Long, G. L. (1989), *Anal. Biochem.* 180, 147-151. The method utilizes 4 primers, 2 15 external (A and D) and 2 internal primers (B and C) that are mutagenic. The strategy involved PCR amplification of a cassette having defined unique restriction enzyme sites surrounding the desired mutation. This cassette was subcloned into a wild-type FLINT vector backbone.
- 20 Following PCR amplification, inserts were cloned into commercial PCR cloning vector pCR2.1TOPO (Invitrogen) and submitted for DNA sequencing prior to subcloning into an expression vector.

- As an example, the cassette for construction of the 25 analog Ala12 to Asn, comprising NheI and KpnI ends, was generated using the following oligonucleotide primers:

CF119 (A): gag cta gcc acc atg agg gcg ctg gag ggg cca  
ggc ctg tcg ctg

CF120(B): GTC TCG TTG TCC CGC CAT GGG TAG GTG GGT GTT  
TCT GCC ACT CCG CGT ACA G

CF121(C): ggc aga aac acc cac cta ccc atg gcg gga caa  
cga gac agg gga gcg gct g

CF122(D): GTC GAT GAC GGC ACG CTC ACA CTC CTC AGC TCC  
TGG TAC CCT GGT GCT G

By increasing the sialic acid content of the FLINT molecule, analogs of the present invention have an increased clearance time from serum. Analogs having greater sialic  
5 acid content than native human FLINT are generated by adding glycosylation sites which do not perturb the secondary or tertiary conformation required for biological activity. Preferably, an analog of FLINT has 1, 2 or 3 additional sites for N-glycosylation. Each additional N-linked site can  
10 provide up to four additional sialic acids per molecule of FLINT analog.

According to the present invention, FLINT isoforms can be separated by a variety of techniques including isoelectric focusing (IEF). When placed in a pH gradient and  
15 subjected to an electric field, proteins will migrate to the point at which they have no net charge. This is the isoelectric point (pI) of the protein. Each distinct band observed on IEF represents molecules that have a particular pI and therefore the same overall charge, termed an isoform.  
20 The term "FLINT isoform" as used herein refers to FLINT preparations having about the same pI, as measured by any suitable technique, and the same amino acid sequence.

Other means for separating FLINT isoforms include fractionation over an ion exchange column. Preferably  
25 isoforms are separated by liquid chromatography (LC).

In a preferred embodiment, FLINT analogs are the product of the expression of an exogenous DNA sequence transfected into a non-human eucaryotic host cell (i.e. "recombinant FLINT"). A recombinant FLINT analog is  
5 advantageously produced and purified according to the procedures described in commonly owned PCT applications WO 00/58466, WO 00/58465, and WO 99/50413, hereby incorporated by reference. For example, the native FLINT cDNA was incorporated into vector pcDNA3 DHFR which provides the CMV  
10 promoter to drive FLINT gene expression and DHFR selection. DG44-C.B4 CHO cells were transfected with linearized vector by electroporation. For non-selective propagation, cells were grown in Ex-Cell 302 Medium (JRH BioSciences), 1X HT Supplement (GibcoBRL), 1X dextran sulfate (Sigma) and 6 mM  
15 L-glutamine (GibcoBRL). For selective growth, cells were placed in Ex-Cell 302 Medium (JRH BioSciences), 1X HT Supplement (GibcoBRL), 1X dextran sulfate (Sigma), 6 mM L-glutamine (GibcoBRL), and Methotrexate (20 mM stock, USP).

After electroporation cells were placed in non-  
20 selective growth medium to recover for 72 hours. Plating was done using 96 well dishes. The cells were plated at various cell densities, and under various levels of methotrexate (MTX) selective pressure. When colony formation was visible the plates were screen by ELISA. The wells were moved into  
25 24 well dishes and expanded to generate enough cells for a full expression study evaluation.

Master wells were amplified at different levels of methotrexate. All master wells showed increased levels of expression at the end of the amplification step. Two master  
30 wells were cloned using FACS cloning.

Discrete isoforms of a given recombinant FLINT analog are contemplated herein corresponding to FLINT molecules

having from 0, 1, 2, 3, 4, 5, 6, 7, 8 or greater number of sialic acids per molecule of analog. Increasing the number of sialic acid residues per molecule of FLINT has the effect of slowing the clearance of FLINT *in vivo*.

5       As demonstrated herein, the *in vivo* clearance of FLINT and FLINT analog variants correlates with sialic acid content. (See Examples 3 and 4). Specifically, FLINT analogs having greater sialic acid content, as expressed, for  
10       example, by the average sialic acid content, are cleared more slowly than the corresponding analog having lesser sialic acid content. Native FLINT possesses one N-linked site and one O-linked site, whereas the analogs possess at least one additional N-linked site and/or O-linked sites.

#### 15       FLINT Analog Isoforms

          The subject invention provides compositions of FLINT analog isoforms. The specific isoforms of FLINT obtained in accordance with the present invention, and their properties, may vary depending upon the source of the starting material.  
20       In a preferred embodiment, the invention relates to a FLINT isoform composition having an average sialic acid content of about 0.5 sialic acid residues per molecule of FLINT analog; alternatively, an average of about 1.0 sialic acids per molecule of FLINT analog; alternatively, an average of about  
25       1.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 2.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 2.5 sialic acids per molecule of FLINT analog;  
30       alternatively, an average of about 3.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 3.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 4.0 sialic acids per

molecule of FLINT analog; alternatively, an average of about 4.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 5.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 5.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 6.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 6.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 7.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 7.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 8.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 8.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 9.0 sialic acids per molecule of FLINT analog; alternatively, an average greater than about 9.0 sialic acids per molecule of FLINT analog.

The compositions of the invention, defined in terms of average sialic acid content, may comprise different FLINT analog isoforms, in other words, a composition of heterogeneous isoforms. For example, an N-linked site theoretically can accommodate 0, 1, 2, 3, or 4 sialic acids per molecule of analog while an O-linked site can theoretically accommodate 0, 1, or 2 sialic acids per molecule of analog. Thus, a composition having an average sialic acid content of about 1 could comprise multiple sialic acid isoforms. For example, each molecule of analog could independently have 1 sialic acid at each N-linked site or 1 sialic acid at the O-linked site, to give an average sialic acid content of 1. Alternatively, molecules with more than 1 sialic acid per molecule of analog could comprise the composition, so long as the average sialic acid content of 1

was maintained. Compositions of the invention comprise any combination of such isoforms that fall within a particular scope of the average sialic acid content. Thus, in one embodiment the compositions of the invention comprise a mixture of isoforms having an average number of sialic acids per FLINT analog molecule.

The invention also provides methods for preparing FLINT analog isoform compositions. These methods include isolation by techniques such as preparative isoelectric focusing, ion exchange chromatography, chromatofocusing.

In general, ion exchange chromatography and chromatofocusing involve application of either conditioned medium containing FLINT analog, or purified material, to a column resin under conditions that permit binding of some or all of the FLINT analog isoforms to the resin. It is preferable to apply the protein to the column at about pH 5. After washing the column with buffer at about pH 5, FLINT analog isoforms that remain bound on the ion exchange column are eluted by increasing the salt concentration of the buffer. For chromatofocusing, isoforms are eluted from the column by a gradient of decreasing pH, or by washing the column with a high concentration of salt.

FLINT analog molecules have N-linked or O-linked oligosaccharide structures which may limit the sialic acid content of the molecule. For example, tetra-antennary (four-branched) N-linked oligosaccharides provide four possible sites for sialic acid attachment, while bi- and triantennary oligosaccharide chains, which can substitute for the tetra-antennary form at asparagine-linked sites, commonly have only two or three sialic acids attached. O-linked oligosaccharides commonly provide two sites for sialic acid attachment.

Thus, native FLINT molecules can accommodate a total of 8 sialic acid residues provided the single N-linked oligosaccharides is tetra-antennary. Analogs would provide 0, 1, 2, 3, or 4 additional sialic acids per additional N-linked site, and 0, 1, or 2 additional sialic acids per additional O-linked site. The N-linked oligosaccharides of FLINT contain sialic acid in both an  $\alpha$ -2,3 and an  $\alpha$ -2,6 linkage to galactose (Takeuchi et al. J. Biol. Chem. 263, 3657(1988)). Typically the sialic acid in the  $\alpha$ -2,3 linkage is added to galactose on the mannose  $\alpha$ -1,6 branch, and the sialic acid in the  $\alpha$ -2,6 linkage is added to the galactose on the mannose  $\alpha$ -1,3 branch. The enzymes that add these sialic acids ( $\beta$ -galactoside  $\alpha$ -2,3 sialyltransferase and  $\beta$ -galactoside  $\alpha$ -2,6 sialyltransferase) are most efficient at adding sialic acid to the mannose  $\alpha$ -1,6 and mannose  $\alpha$ -1,3 branches respectively.

Mammalian cell cultures may be screened for cells that preferentially add tetra-antennary chains to recombinant FLINT analog, thereby maximizing the number of sites for sialic acid attachment. Dihydrofolate reductase (DHFR) deficient Chinese Hamster Ovary (CHO) cells are commonly used for the production of recombinant glycoproteins including recombinant FLINT. These cells do not express the enzyme  $\beta$ -galactoside  $\alpha$ -2,6 sialyltransferase, and therefore do not add sialic acid in the  $\alpha$ -2,6 linkage to N-linked oligosaccharides of glycoproteins produced in these cells. (Mutsaers et al. Eur. J. Biochem. 156, 651 (1986); Takeuchi et al. J. Chromatogr. 400, 207 (1987)). Consequently, recombinant FLINT produced in CHO cells lacks sialic acid in the 2,6 linkage to galactose (Sasaki et al. (1987), supra; Takeuchi et al. (1987), supra). Therefore, in one embodiment



of the invention, FLINT analog isoforms are made in CHO cells that are transfected with a functional  $\beta$ -galactoside  $\alpha$ -2,6 sialyltransferase gene to give incorporation of sialic acid in  $\alpha$ -2,6 linkage to galactose. See Lee et al. J. Biol. Chem. 264, 13848 (1989), hereby incorporated by reference, for a disclosure of techniques for creating modified CHO cells, or other mammalian host cells.

#### Enzymatically Enhanced Sialylation of FLINT

Also contemplated by the present invention is a method for enhancing the sialylation of FLINT analog isoforms by enzymatic modification *in vitro*.

The circulatory lifetime of glycoproteins such as FLINT analog in blood is highly dependent on the composition and structure of N-linked oligosaccharides. In general, maximal plasma half-life of a glycoprotein requires that its N-linked carbohydrate groups terminate in the sequence NeuAcGalGlcNAc. Without a terminal sialic acid residue (NeuAc), a glycoprotein is rapidly cleared from the blood by receptors that recognize the exposed Gal residues. For this reason, ensuring high sialylation of therapeutic proteins such as FLINT analog is important for commercial development.

Although much is known about the complexity of carbohydrate structures on glycoproteins, attempts to specify post-translational glycosylation in cultured cells have not kept pace with advances in technology for gene expression, and therefore, incomplete glycosylation of secreted recombinant glycoproteins, including FLINT analogs, is common. One solution to this problem is to use isolated

glycosyltransferases to complete carbohydrate chains *in vitro*.

Optimal glycosylation may be difficult to achieve using mammalian cell culture systems. Under conditions of large  
5 scale growth, overproduction of a protein backbone comprising a glycoprotein can exceed the host cells capacity to achieve full sialylation.

A method of the present invention comprises use of sialyltransferase to add sialic acids to an acceptor site(s)  
10 on FLINT analog, preferably said site(s) having a galactosyl unit. The method for enhancing sialylation of FLINT follows that disclosed in U.S. Patent No. 6,030,815, herein incorporated by reference. Essentially, the method comprises the steps of adding sialyltransferase to a sample of FLINT  
15 analog and a catalytic amount of a CMP-sialic acid synthetase, a sialic acid, CTP, and a soluble divalent metal cation, including  $Mn^{+2}$ ,  $Mg^{+2}$ ,  $Ca^{+2}$ ,  $Co^{+2}$ , and  $Zn^{+2}$ . Preferably, the divalent ion concentration is maintained between 2mM and 75 mM. Alternatively, the reaction may further comprise a  
20 CMP-sialic acid recycling system, as disclosed in U.S. Patent 6,030,815. A commercially available system for carrying out the reactions, GlycoAdvance®, is available from Neose Technologies, Inc. (Horsham, PA).

In one embodiment of the invention, FLINT analog is  
25 produced recombinantly in a suitable mammalian cell type, for example, CHO cells, by transfection with a suitable vector for expressing FLINT analog. Culture supernatants containing FLINT analog are concentrated and processed using the GlycoAdvance® system, or comparable commercial or non-  
30 commercial system. FLINT analog having enhanced sialylation is recovered using standard purification techniques.

Also comprehended by the invention are pharmaceutical compositions comprising a therapeutically effective amount of a FLINT analog isoform, having an average sialic acid content of about 0.5 sialic acid residues per molecule of FLINT analog; alternatively, an average of about 1.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 1.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 2.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 2.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 3.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 3.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 4.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 4.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 5.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 5.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 6.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 6.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 7.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 7.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 8.0 sialic acids per molecule of FLINT analog; alternatively, an average of about 8.5 sialic acids per molecule of FLINT analog; alternatively, an average of about 9.0 sialic acids per molecule of FLINT analog; alternatively, an average greater than about 9.0 sialic acids per molecule of FLINT analog, together with a suitable diluent, adjuvant and/or carrier

useful in therapeutic applications. A therapeutically effective amount as used herein refers to that amount which provides therapeutic effect for a given condition and administration regimen. The administration of FLINT analog isoforms is preferably by the intravenous route.

#### Therapeutic Applications

The clinical utility for the FLINT analog isoforms of the invention is expected to be substantial. Active FLINT analog isoforms inhibit the binding of Fas to FasL and LIGHT to LTBR and TR2/HVEM receptors, and can be used to treat or prevent a disease and/or condition that may be associated with such binding.

Many diseases and/or conditions involving FasL/Fas are potentially amenable to therapy with FLINT analog isoforms. Examples of suitable diseases and/or conditions include the following.

Inflammatory/autoimmune diseases - Rheumatoid arthritis, inflammatory bowel disease, graft-versus-host disease, insulin-dependent diabetes, SIRS/sepsis/MODS, pancreatitis, psoriasis, multiple sclerosis, Hashimoto's thyroiditis, Grave's disease, transplant rejection, SLE, autoimmune gastritis, fibrosing lung disease.

Infectious diseases - HIV-induced lymphopenia, fulminant viral hepatitis B/C, chronic hepatitis/cirrhosis, H. pylori-associated ulceration.

Ischemia/Re-perfusion conditions - Acute coronary syndrome, acute myocardial infarction, congestive heart failure, atherosclerosis, acute cerebral ischemia/infarction, brain/spinal cord trauma, organ preservation during transplant

Other treatments include cytoprotection during cancer treatment, adjuvant to chemotherapy, Alzheimer's, chronic glomerulonephritis, osteoporosis, TTP/HUS, aplastic anemia, myelodysplasia. Also of interest are treatment and  
5 prevention of acute lung injury (ALI)/acute respiratory distress syndrome (ARDS); Ulcerative colitis; and Crohn's disease.

Other diseases for which FLINT analog isoforms are therapeutically useful include rheumatoid arthritis (Elliott  
10 et al., *Lancet* 344:1105-10 (1994)), fibroproliferative lung disease, fibrotic lung disease, HIV (Dockrell et al., *J. Clin. Invest.* 101:2394-2405 (1998)), Ischemia (Sakurai et al. 1998 *Brain Res* 797:23-28), Brain trauma/injury (Ertel et al. 1997 *J Neuroimmunol* 80:93-6), chronic renal  
15 failure (Schelling et al. 1998 *Lab Invest* 78:813-824), Graft-vs-Host Disease (GVHD) (Hattori et al. 1998 *Blood* 11:4051-4055), Cutaneous inflammation (Orteu et al. 1998 *J Immunol* 161:1619-1629), Vascular leak syndrome (Rafi et al. 1998 *J Immunol* 161:3077-3086), *Helicobacter pylori*  
20 infection (Rudi et al. 1998 *J Clin Invest* 102:1506-1514), Goiter (Tamura et al. 1998 *Endocrinology* 139:3646-3653), Atherosclerosis (Sata and Walsh, 1998 *J Clin Invest* 102:1682-1689), IDDM (Itoh et al. 1997 *J Exp Med* 186:613-618), Osteoporosis (Jilka et al. 1998 *J Bone Min Res*  
25 13:793-802), Crohn's Disease (van Dullemen et al. 1995 *Gastroenterology* 109:129-35), organ preservation and transplant (graft) rejection (Lau et al. 1996 *Science* 273:109-112), Sepsis (Faist and Kim. 1998 *New Horizons* 6:S97-102), Pancreatitis (Neoptolemos et al. 1998 *Gut*  
30 42:886-91), Cancer (melanoma, colon and esophageal) (Bennett et al. 1998 *J Immunol* 160:5669-5675), Autoimmune disease (IBD, psoriasis, Down's Syndrome (Seidi et al., *Neuroscience*

Lett. 260:9 (1999), multiple sclerosis (D'Souza et al. 1996 J Exp Med 184:2361-70), Alzheimer's Disease; End-stage renal disease (ESRD); mononucleosis; EBV; Herpes; antibody dependent cytotoxicity; hemolytic and hypercoagulation disorders such as vascular bleeds, DIC (disseminated intravascular coagulation), eclampsia, HELLP (preeclampsia complicated by thrombocytopenia, hemolysis and disturbed liver function), HITS (heparin induced thrombocytopenia), HUS (hemolytic uremic syndrome), and preeclampsia; hematopoietic disorders such as aplastic anemia, thrombocytopenia (TTP) and myelodysplasia; and hemolytic fever caused, for example, by E.bola.

In the case of organ preservation in preparation for harvesting, for instance, a FLINT analog isoform is useful prophylactically to prevent the apoptosis associated with ischemia reperfusion injury to the organ once it is removed from the donor. Suitable media for this purpose are known, for example, the media disclosed in EP 0356367 A2. The method may also include treating the transplant recipient with FLINT analog isoform prior to and/or after the transplant surgery.

There is evidence that ARDS may be mediated by soluble FasL/Fas interaction in humans (Matute-Bello et al., J. Immunol. 163, 2217-2225, 1999). FLINT, by binding to FasL, could inhibit FasL-mediated apoptosis of pneumocytes and/or endothelial cells, thus inhibiting or preventing the progression from acute inflammatory insult to ALI, and from ALI to ARDS.

Therefore, in another embodiment, the present invention relates to the use of a FLINT analog isoforms to inhibit and/or treat ALI and/or ARDS comprising the administration

of a therapeutically effective amount of analog isoform to a person in need thereof.

In another embodiment, the present invention relates to the use of a FLINT analog isoform to treat and/or inhibit  
5 chronic obstructive pulmonary disease (COPD) in a patient in need thereof by administering a therapeutically effective amount of FLINT analog isoform.

In another embodiment the present invention relates to the use of a FLINT analog isoform to inhibit and/or treat  
10 pulmonary fibrosis (PF). For example, FLINT analog isoform can be administered acutely at the time of an inflammatory insult to the lung (e.g. during bleomycin treatment) to prevent PF from occurring.

A "subject" is a mammal in need of treatment, preferably  
15 a human, but can also be an animal in need of veterinary treatment, e.g., domestic animals (e.g., dogs, cats, and the like), farm animals (e.g., cows, sheep, pigs, horses, and the like) and laboratory animals (e.g., rats, mice, guinea pigs, and the like).

20 An "effective amount" of FLINT analog isoform is an amount which results in a sufficient inhibition of one or more processes mediated by the binding of Fas to FasL or LIGHT to LTBR and/or TR2/HVEM so as to achieve a desired therapeutic or prophylactic effect in a subject with a  
25 disease or condition that may be associated with aberrant Fas/FasL binding and/or LIGHT mediated binding.

Alternatively, an "effective amount" of FLINT analog isoform is a quantity sufficient to achieve a desired therapeutic and/or prophylactic effect in a subject with inflammation  
30 caused by FasL-induced neutrophil activation or any of the other aforementioned diseases associated with aberrant FasL activity.

A "desired therapeutic and/or prophylactic effect" in a subject with a disease or condition includes the amelioration of symptoms, or delay in onset of symptoms, associated with such disease. Alternatively, a "desired  
5 therapeutic and/or prophylactic effect" includes an increased survival rate or increased longevity for the subject with the disease.

The amount of FLINT analog isoform administered to the individual will depend on the type and severity of the  
10 disease and on the characteristics of the individual, such as general health, age, sex, body weight and tolerance to drugs. It will also depend on the degree, severity and type of disease. The skilled artisan will be able to determine appropriate dosages depending on these and other factors.

15 As a general proposition, the total pharmaceutically effective amount of the FLINT analog isoform molecules of the present invention administered parenterally per dose will be in the range of about 1  $\mu\text{g/kg/day}$  to 10  $\text{mg/kg/day}$  of patient body weight, particularly 2  $\text{mg/kg/day}$  to 8  
20  $\text{mg/kg/day}$ , more particularly 2  $\text{mg/kg/day}$  to 4  $\text{mg/kg/day}$ , even more particularly 2.2  $\text{mg/kg/day}$  to 3.3  $\text{mg/kg/day}$ , and finally 2.5  $\text{mg/kg/day}$ , although, as noted above, this will be subject to therapeutic discretion. More preferably, this dose is at least 0.01  $\text{mg/kg/day}$ . If given continuously, a  
25 FLINT analog isoform of the present invention is typically administered at a dose rate of about 1  $\mu\text{g/kg/hour}$  to about 50  $\mu\text{g/kg/hour}$ , either by 1-4 injections per day or by continuous subcutaneous infusions, for example, using a mini-pump. An intravenous bag solution may also be  
30 employed. The length of treatment needed to observe changes and the interval following treatment for responses to occur appears to vary depending on the desired effect.



Pharmaceutical compositions comprising FLINT analog isoform(s) of the present invention may be administered orally, rectally, intracranially, parenterally, intracisternally, intravaginally, intraperitoneally, 5 topically (as by powders, ointments, drops or transdermal patch), transdermally, intrathecally, buccally, or as an oral or nasal spray. By "pharmaceutically acceptable carrier" is meant a non-toxic solid, semisolid or liquid filler, diluent, encapsulating material or formulation auxiliary of any type. The term "parenteral" as used herein includes, 10 but is not limited to, modes of administration which include intravenous, intramuscular, intraperitoneal, intrasternal, subcutaneous and intraarticular injection, infusion and implants comprising FLINT analog isoforms.

15 The FLINT analog isoforms of the present invention are also suitably administered by sustained-release systems. Suitable examples of sustained-release compositions include semi-permeable polymer matrices in the form of shaped articles, e.g., films, or microcapsules. Sustained-release 20 matrices include polylactides (U.S. Pat. No. 3,773,919, EP 58,481), copolymers of L-glutamic acid and gamma-ethyl-L-glutamate (Sidman, U. et al., *Biopolymers* 22:547-556 (1983)), poly (2-hydroxyethyl methacrylate) (R.Langer et al., *J. Biomed. Mater. Res.* 15:167-277 (1981), and R. 25 Langer, *Chem. Tech.* 12:98-105 (1982)), ethylene vinyl acetate (R. Langer et al., *Id.*) or poly-D-(-)-3-hydroxybutyric acid (EP 133,988). Other sustained-release compositions also include liposomally entrapped FLINT analog isoform. Such liposomes are prepared by methods known per 30 se: DE 3,218,121; Epstein et al., *Proc. Natl. Acad. Sci. (USA)* 82:3688-3692 (1985); Hwang et al., *Proc. Natl. Acad. Sci. (USA)* 77:4030-4034 (1980); EP 52,322; EP 36,676; EP

88,046; EDP 143,949; EP 142,641; Japanese Pat. Appl. 83-118008; U.S. Pat. Nos. 4,485,045 and 4,544,545; and EP 102,324. Ordinarily, the liposomes are of the small (about 200-800 Angstroms) unilamellar type in which the lipid  
5 content is greater than about 30 mol. percent cholesterol, the selected proportion being adjusted for the optimal TNFR polypeptide therapy.

For parenteral administration, the FLINT analog isoforms of the present invention are formulated generally  
10 by mixing at the desired degree of purity, in a unit dosage injectable form (solution, suspension, or emulsion), with a pharmaceutically acceptable carrier, i.e., one that is non-toxic to recipients at the dosages and concentrations employed and is compatible with other ingredients of the  
15 formulation. For example, the formulation preferably does not include oxidizing agents and other compounds that are known to be deleterious to polypeptides.

The FLINT analog isoforms of the present invention are typically formulated in suitable vehicles at a concentration  
20 of about 0.1 mg/ml to 100 mg/ml, preferably 1-10 mg/ml, at a pH of about 3 to 8. It will be understood that the use of certain excipients, carriers, or stabilizers will result in the formation of salts of the FLINT analog molecules of the present invention.

25 The invention also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the pharmaceutical compositions of the invention. Associated with such container(s) can be a notice in the form prescribed by a governmental agency  
30 regulating the manufacture, use or sale of pharmaceuticals or biological products, which notice reflects approval by the agency of manufacture, use or sale for human

administration. In addition, the FLINT analog isoforms of the present invention may be employed in conjunction with other therapeutic compounds.

The following examples are offered to more fully illustrate the invention, but are not to be construed as limiting the scope thereof.

#### EXAMPLE 1

##### Isolation of Recombinant FLINT Analog A12N Isoforms

10 A bicistronic expression vector was constructed by inserting into mammalian expression vector pGTD (Gerlitz, B. et al., 1993, Biochemical Journal 295:131) a PCR fragment encoding an "internal ribosome entry site"/enhanced green fluorescent polypeptide (IRES/eGFP). The new vector, designated pIG3, contains the following elements: the E1a-responsive GBMT promoter (D. T. Berg et al., 1993 BioTechniques 14:972; D.T. Berg et al., 1992 Nucleic Acids Research 20:5485); a multiple cloning site (MCS); the IRES sequence from encephalomyocarditis virus (EMCV); the eGFP coding sequence (Cormack, et al., 1996 Gene 173:33, Clontech); the SV40 small "t" antigen splice site/polyadenylation sequences; the SV40 early promoter and origin of replication; the murine dihydrofolate reductase (dhfr) coding sequence; and the ampicillin resistance gene and origin of replication from pBR322.

Based on the human FLINT cDNA sequence (e.g. SEQ ID NO:3), forward and reverse PCR primers were synthesized bearing *BclI* restriction sites at their respective 5' ends. These primers were used to amplify the FLINT cDNA. The FLINT cDNA orientation and nucleotide sequence was confirmed by restriction digest and double stranded sequencing of the insert. The approximately 900 base pair amplified FLINT

analog PCR product was digested with restriction endonucleases NheI and XbaI, respectively, to generate a fragment bearing NheI and XbaI sticky ends. This fragment was subsequently ligated into a unique XbaI site of pIG3 to  
5 generate recombinant plasmid pIG3-FLINT.

The recombinant pIG3-FLINT plasmid carries the FLINT gene and encodes resistance to methotrexate. In vitro mutagenesis processes were followed to change Ala 12 to Asn, in order to introduce an additional N-linked glycosylation  
10 site as described elsewhere in this disclosure. AV12 RGT18 cells are transfected using a calcium phosphate procedure with the recombinant pIG vector. Cells resistant to 250 nM methotrexate are selected and pooled. The pool of resistant clones is subjected to fluorescence-assisted cell sorting  
15 (FACS), and cells having fluorescence values in the top 5% are sorted into a pool, and as single cells. High fluorescence pools are subjected to two successive sorting cycles. Pools and individual clones from the first and second cycles are analyzed for FLINT analog production by  
20 ELISA. Pools or clones expressing FLINT analog at the highest level are used for scale-up and FLINT analog purification.

Large-scale production of FLINT analog A12N is carried out by growing stable clones of AV12 RGT 18 cells  
25 transfected with an expression vector that expresses A12N. After reaching confluence, cells are further incubated for 2-3 more days to secrete maximum amount of FLINT analog into the growth medium. Medium containing FLINT analog is adjusted to 0.1 % CHAPS and concentrated in an Amicon  
30 ProFlux M12 tangential filtration system to 350 ml. The concentrated medium is adjusted to pH 6.0 and passed over a SP Sepharose Fast Flow (Pharmacia, 500 ml) at a flow rate of

7 ml/min. The column is washed with buffer A (20 mM MOPS, 0.1 % CHAPS, pH 6.0) until the absorbance (280 nm) returns to baseline and bound polypeptides are eluted with a linear gradient from 0 to 1 M NaCl (in buffer A) developed over  
5 four column volumes. Fractions containing FLINT only are pooled and passed over Vydac C4 column (100 ml) equilibrated with 0.1 % TFA/H<sub>2</sub>O at a flow rate of 10 ml/min. This material is passed over a 16/60 Superdex 200 sizing column (Pharmacia) equilibrated with PBS, 0.5 M NaCl, pH 7.4.  
10 Fractions containing FLINT analog are analyzed by SDS-PAGE, and the N-terminal sequence of the purified polypeptide confirmed to be FLINT.

In an alternative purification scheme, concentrated and clarified medium containing FLINT analog is passed over Blue  
15 Sepharose DAC and eluted with 7M urea, 1M NaCl, pH 8. Eluted material is further purified on CG71 reverse phase chromatography and eluted with 35% acetonitrile, 0.6 M NaCl, pH 7.4. The eluted material is purified further on SP650M cation exchange with elution in 30% acetonitrile, pH 2.5..  
20 Following this step, eluted material is subjected to solvent exchange by TFF into buffer for bulk freezer storage.

FLINT A12N has two potential N-linked glycosylation sites, at Asn 12 and Asn144 of SEQ ID NO:1. To characterize the oligosaccharide structure of A12N produced in AV12  
25 cells, samples of intact FLINT analog A12N, and FLINT A12N that had been treated with neuraminidase and HEXase II to release terminal GalNAc and GlcNAc residues (desialylated FLINT), is analyzed by capillary HPLC/ESI-MS. The oligosaccharides released by this treatment are labeled with  
30 2-aminobenzamide and analyzed by weak anion exchange (WAE) HPLC and by LC mass spectrophotometry (LC/MS).

## EXAMPLE 2

Oligosaccharide Profile of FLINT A12N by HPLC/ESI-MS  
and Fluorescence HPLC

Recombinant A12N FLINT is expressed in AV12 cells and  
5 purified as in Example 1. Intact FLINT A12N is directly  
analyzed by a capillary HPLC/ESI-MS, or treated by  
neuraminidase, or HEXase II, followed by HPLC/ESI-MS. The  
oligosaccharide structures are calculated based on the  
obtained masses and the expected FLINT analog peptide  
10 backbone mass. Fluorescence labeled FLINT analog  
oligosaccharides are fractionated by weak anion exchange  
(WAE) HPLC. The fractions are collected and identified by  
liquid chromatography/mass spectrophotometry (LC/MS).  
Neuraminidase, HEXase II Treatment of A12N Ten microliters  
15 of a solution containing FLINT A12N (~0.43 mg/ml in PBS,  
0.5M NaCl) is mixed with 8 uL of 50 mM NaOAc buffer, pH 5.2,  
and 2 uL of neuraminidase solution (1 unit/mL). The mixture  
is incubated at 37°C for 2 hours. Seven microliters of the  
mixture is used for capillary HPLC/MS analysis and two  
20 microliters of HEXase II enzyme solution (Glyko, Inc.) is  
added to the remaining solution, which is incubated at 37°C  
for 3 hours before HPLC/MS analysis.  
Weak Anion Exchange (WAE) HPLC of Fluorescence labeled  
Oligosaccharides. A 200 ul Aliquot of thawed FLINT A12N  
25 solution containing approximately 0.2 mg of protein is mixed  
with 60 mg urea, 17.6 ul of 3 M Tris buffer (pH 8.0) and 3  
ul of 50 mg/mL dithiothreitol and the mixture was incubated  
at 37°C for 10 min. The sample is alkylated by adding 5 ul  
of 100 mg/mL iodoacetic acid solution and incubating at  
30 ambient temperature in the dark for 10 min. Samples are  
desalted on a disposable gel filtration column and  
oligosaccharides released by treatment with 1 unit N-

glycosidase F solution at 37°C for 2 hours. The deglycosylated protein is precipitated by adjusting pH with 10% (v/v) acetic acid solution. A 300 ul aliquot of oligosaccharide solution is dried and labeled with 2-aminobenzamide dye. After the excess dye is removed using a P-2 spin column, the labeled oligosaccharide solution is analyzed by WAE-HPLC with a fluorescence detector.

Desialylation of WAE HPLC fractions of 2-AB labeled FLINT

Oligosaccharides. Each of the collected WAE HPLC fractions is transferred into two vials and dried using a centrifugation vacuum system. Fifteen microliters (containing 2.5 munits neuraminidase) of 15 mM NaOAc buffer, pH 5.2 is added to one vial for each fraction. Five microliters of the mixture is used for LC/MS analysis after incubation at room temperature for 10 to 15 hours. For intact oligosaccharide fraction, fifteen microliters of H<sub>2</sub>O is added into each vial and 5 ul of the solution is used for LC/MS analysis.

Capillary HPLC/ESI-MS. A Beckman System Gold equipped with a Model 126 solvent delivery module is used. The HPLC buffer (A: 0.15% formic acid in H<sub>2</sub>O and B: 0.12% formic acid in ACN) is pumped through a T split, in which a Zorbax 300SB C18, 2.1x150 mm column is attached on one exit and a manual injection valve, Vydac capillary column (C18, 0.3x150 mm) and a API UV detector (785A) at the other exit. The HPLC stream from the capillary directly passes to the mass spectrometer through a fused silica transfer line. Beckman solvent deliver system is pumped at 0.2 ml/min with the following gradient

Time (min)	0	2	42	43	45	46	57
------------	---	---	----	----	----	----	----

Buffer B%            10    10    45    90    90    10    10

After the split, about 5-6 ul/min of HPLC stream is diverted to the capillary column. About 2 ug FLINT A12N is injected to capillary column per run. API UV detector is set lambda = 214 nm and the data stored in HP1000. A PE Sciex API III mass spectrometer equipped with an articulated ionspray source is used in these studies under the condition of CC 1, OR 55 V and ISV 4800 V, Q1 scan from 900 to 1500 or 1000 to 1400, 0.33 or 0.20/step, 2 to 3 ms dwell time and 6 sec/scan. For oligosaccharide fraction analysis, a short solvent gradient program is used.

FLINT A12N has two potential N-glycosylation sites. The theoretical molecular weight of the protein alone is 29736.9 Da. The mass spectrum and reconstructed mass spectrum of FLINT A12N are analyzed for the molecular weights of the primary signals.

Based on these experimental results, the primary A12N glycoforms are assigned.

The fluorescence-labeled FLINT A12N oligosaccharides are fractionated into multiple peaks on WAE HPLC according to their negative charge number. The primary structures of oligosaccharides are identified in each fraction.

Because FLINT A12N contains two glycosylation sites, sialic acid content can be calculated based on the results of WEA HPLC as follows:

Sialic Acid Content =  $\sum$  sialylation degree of fraction  $\times$  percentage of fraction/100

Sialic acid content may also be determined by a modification of the procedure of Jourdian et al. J. Biol. Chem. 246, 430 (1971). The sialic acid residues are cleaved from glycoproteins by hydrolysis with 0.35M sulfuric acid at



80° C for 30 minutes and the solutions neutralized with sodium hydroxide prior to analysis. In order to estimate the amount of protein present, a Bradford protein assay (Bradford Anal. Biochem. 72, 248 (1976)) using recombinant FLINT as standard is performed using the assay reagents and the micro-method procedure supplied by Bio-Rad.

### EXAMPLE 3

#### Effect of Higher Sialylation on Clearance of A12N FLINT

10       The pharmacokinetics of different lots of FLINT A12N are examined for the influence of sialic acid content on the clearance in male Cynomolgus monkeys. Sialic acid content may be determined as in Example 2, based on LC/MS and oligosaccharide profiling.

15       Lots of A12N are administered as a single intravenous bolus dose (0.5 mg/kg) and blood samples obtained over a 48 h period after dosing.

      Plasma samples from treated animals are analyzed for concentrations of FLINT(A12N) using a sandwich ELISA method  
20   employing affinity purified rabbit polyclonal anti-FLINT antibodies.

      After intravenous administration, lots of A12N are  
~   analyzed for clearance from the plasma of treated animals. The data indicate that the extent of terminal sialylation on  
25   the carbohydrate moieties present on FLINT A12N have an influence on the clearance kinetics of the compound after administration by the intravenous route. The increased rate of clearance of poorly sialylated molecules from the  
      circulation is most likely mediated through hepatic  
30   asialoglycoprotein receptors via exposed terminal galactose residues.

## EXAMPLE 4

Effect of Higher Sialylation on Clearance of FLINT Analogs

The pharmacokinetics of multiple lots FLINT A12N are examined for the influence of sialic acid content on  
5 clearance in mammals, for example, male Cynomolgus monkeys. Oligosaccharide and sialic acid profiling using LC-MS is used to ascertain sialylation differences between lots.

Lots of FLINT A12N having differing average sialic acid content are administered as a single intravenous bolus dose  
10 to Cynomolgus monkeys (0.5 mg/kg), and blood samples are obtained over a 48 hour period after dosing.

Plasma samples from treated animals are analyzed for concentrations of FLINT A12N using a sandwich ELISA method employing affinity purified rabbit polyclonal anti-FLINT  
15 antibodies. The capture antibody recognizes the N-terminal portion of FLINT. The sandwich antibody is a biotinylated polyclonal antibody which recognizes the C-terminal portion of FLINT.

The effect of sialylation is determined by measuring  
20 the clearance kinetics of the A12N analog after administration by the intravenous route.

## EXAMPLE 5

Fractionation of Recombinant FLINT Analog Isoforms Using a  
25 Low pH Gradient on SP-Sepharose

FLINT analog isoforms are separated using a gradient of decreasing pH and increasing ionic strength. For example, concentrated and filtered FLINT A12N-containing medium prepared as in Example 1 is loaded onto a column of SP-  
30 Sepharose at a ratio of approximately 20 mg total protein/mL gel. The column is then washed with approximately three column volumes of 20 mM MOPS, pH 5.5. FLINT A12N isoforms

are eluted from the column using a gradient starting with 20 mM MOPS, pH 5.5 and running to 20 mM MOPS, 600 mM NaCl, pH 5.5. The total volume of the gradient is approximately 40 column volumes.

5

## EXAMPLE 6

Analogues of FLINT Having Additional Glycosylation Sites

FLINT analogues having additional carbohydrate attachment sites are described elsewhere in this specification.

10 Mutations are introduced into the native FLINT cDNA using well known *in vitro* mutagenesis techniques.

FLINT variant A12N was constructed by mutagenic PCR starting from a wild-type FLINT template. See e.g. Saiki R. K. et al. *Science* 239:487-491 (1988), and "Current Protocols in Molecular Biology", Vol 1, section 8.5.7 (John Wiley and Sons, Inc. publishers), the entire contents of which are  
15 herein incorporated by reference.

The mutagenic PCR process involves a "SOEing" reaction (i.e. Strand Overlap Extension) to create specific mutations  
20 in the native FLINT template for the purpose of changing the amino acid sequence at position 12, and further for introducing restriction enzyme tags for identification purposes.

Generally, SOEing reactions require the use of four  
25 primers, two in the forward orientation (termed A, SEQ ID NO:5, and C, SEQ ID NO:7) and two in the reverse orientation (termed B, SEQ ID NO:6 and D, SEQ ID NO:8). The SOEing reaction amplifies a nucleic acid sequence (e.g. gene sequence) in two stages. The first step amplifies a portion  
30 of the gene by performing an A to B reaction followed by a separate C to D reaction. In constructing the A12N mutant, the B and C primers were targeted to the same area of the

gene but on opposite strands. Mismatch priming from both oligonucleotide primers produces the mutation. After these two reactions were completed, the products were isolated and mixed for use as template for the A to D reaction, which  
5 yields the desired mutated product.

The primers involved in the cloning of A12N were:

CF119 (A): gag cta gcc acc atg agg gcg ctg gag ggg cca  
ggc ctg tcg ctg

CF120(B): GTC TCG TTG TCC CGC CAT GGG TAG GTG GGT GTT  
TCT GCC ACT CCG CGT ACA G

CF121(C): ggc aga aac acc cac cta ccc atg gcg gga caa  
cga gac agg gga gcg gct g

CF122(D): GTC GAT GAC GGC ACG CTC ACA CTC CTC AGC TCC  
TGG TAC CCT GGT GCT G

The amplified fragment carrying the A12N mutation was sub-cloned using an 5' NheI site (GAGCTA) and a 3' KpnI site (GAGGAG). The native FLINT sequence has a naturally  
10 occurring internal KpnI site around amino acid position 176. The amplified mutant fragment was incorporated into the full length FLINT sequence as follows:

First, the amplified fragment was placed into an intermediate vector, pCR2.1- TOPO, which utilizes the  
15 adenine overhangs established after PCR for ligation. And second, the mutated fragment was used to replace the corresponding segment in the wild type FLINT gene by directional ligation.

While the invention has been described in what is  
20 considered to be its preferred embodiments, it is not to be limited to the disclosed embodiments, but on the contrary,

is intended to cover various modifications and equivalents included within the spirit and scope of the appended claims, which scope is to be accorded the broadest interpretation so as to encompass all such modifications and equivalents.

We Claim:

1. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a  
5 change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 0.5 sialic acids per molecule of FLINT analog.
2. A composition comprising a FLINT analog isoform, said  
10 analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 1.0 sialic acids per molecule of FLINT analog.
- 15 3. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 1.5 sialic acids per molecule of FLINT  
20 analog.
4. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid  
25 content of about 2.0 sialic acids per molecule of FLINT analog.
5. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation  
30 site, said composition having an average sialic acid content of about 2.5 sialic acids per molecule of FLINT analog.

6. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 3.0 sialic acids per molecule of FLINT analog.
7. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 3.5 sialic acids per molecule of FLINT analog.
8. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 4.0 sialic acids per molecule of FLINT analog.
9. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 4.5 sialic acids per molecule of FLINT analog.
10. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 5.0 sialic acids per molecule of FLINT analog.
11. A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a

change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 5.5 sialic acids per molecule of FLINT analog.

5 12.A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 6.0 sialic acids per molecule of FLINT  
10 analog.

13.A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid  
15 content of about 6.5 sialic acids per molecule of FLINT analog.

14.A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation  
20 site, said composition having an average sialic acid content of about 7.0 sialic acids per molecule of FLINT analog.

15.A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a  
25 change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 7.5 sialic acids per molecule of FLINT analog.

16.A composition comprising a FLINT analog isoform, said  
30 analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid



content of about 8.0 sialic acids per molecule of FLINT analog.

- 17.A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a  
5 change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 8.5 sialic acids per molecule of FLINT analog.
- 18.A composition comprising a FLINT analog isoform, said  
10 analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid content of about 9.0 sialic acids per molecule of FLINT analog.
- 15 19.A composition comprising a FLINT analog isoform, said analog consisting essentially of SEQ ID NO:3 wherein a change in sequence results in an additional glycosylation site, said composition having an average sialic acid  
20 content of about 9.5 sialic acids per molecule of FLINT analog.
- 20.A pharmaceutical composition comprising a therapeutically effective amount of FLINT analog according to claim 2 and a pharmaceutically acceptable diluent, adjuvant or carrier.

## SEQUENCE LISTING

<110> Eli Lilly and Company

<120> FLINT Analog Glycoforms

<130> X-14727

<140>

<141>

<160> 8

<170> PatentIn Ver. 2.0

<210> 1

<211> 271

<212> PRT

<213> Homo sapiens

<400> 1

Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu

1

5

10

15

Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro

20

25

30

Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His

35

40

45

Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val

50

55

60

Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His

65

70

75

80

-2-

Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe  
                                     85                                    90                                    95

Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro  
                                     100                                    105                                    110

Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr  
                                     115                                    120                                    125

Phe Ser Ala Ser Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn  
                                     130                                    135                                    140

Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His  
                                     145                                    150                                    155                                    160

Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val  
                                     165                                    170                                    175

Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe  
                                     180                                    185                                    190

Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu  
                                     195                                    200                                    205

Ala Pro Glu Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu  
                                     210                                    215                                    220

Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp  
                                     225                                    230                                    235                                    240

Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met  
                                     245                                    250                                    255

Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His  
                                     260                                    265                                    270

&lt;210&gt; 2

&lt;211&gt; 813

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 2

```

gtggcagaaa caccacaccta cccctggcgg gacgcagaga caggggagcg gctgggtgtgc 60
gcccagtgcc cccagggcac ctttgtgcag cgcccggtgcc gccgagacag cccacagacg 120
tgtggcccggt gtccaccgcg ccactacacg cagttctgga actacctgga gcgctgccgc 180
tactgcaacg tcctctgcgg ggagcgtgag gaggaggcac gggcttgcca cgccaccac 240
aaccgtgcct gccgctgccg caccggcttc ttgcgcacg ctggtttctg cttggagcac 300
gcatcgtgtc cacctgggtgc cggcgtgatt gcccggggca ccccagcca gaacacgcag 360
tgccagccgt gccccccagg caccttctca gccagcagct ccagctcaga gcagtgccag 420
ccccaccgca actgcacggc cctgggctctg gccctcaatg tgccaggctc ttcctcccat 480
gacacctgtg gcaccagctg cactggcttc cccctcagca ccagggtacc aggagctgag 540
gagtgtgagc gtgccgtcat cgactttgtg gctttccagg acatctccat caagaggctg 600
cagcgggtgc tgcaggccct cgaggccccg gagggctggg gtccgacacc aagggcgggc 660
cgcgcggcct tgcagctgaa gctgcgtcgg cggtcacgg agctcctggg ggcgaggac 720
ggggcgctgc tgggtcggct gctgcaggcg ctgcgcgtgg ccaggatgcc cgggctggag 780
cggagcgtcc gtgagcgctt cctccctgtg cac 813

```

&lt;210&gt; 3

&lt;211&gt; 300

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 3

```

Met Arg Ala Leu Glu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val Leu
  1              5              10              15

Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly Val Ala Glu
      20              25              30

Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val
    35              40              45

Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg
    50              55              60

Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln
    65              70              75              80

```

Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly  
85 90 95

Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His Asn Arg Ala  
100 105 110

Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu  
115 120 125

His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro  
130 135 140

Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala  
145                    150                    155                    160

Ser Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala  
165 170 175

Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu  
180 185 190

Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala  
195 200 205

Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile  
210 215 220

Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu  
225                      230                      235                      240

Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys  
245 250 255

Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu  
260 265 270

Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu  
275 280 285

Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His  
 290 295 300

<210> 4  
 <211> 936  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> CDS  
 <222> (25)..(924)

<400> 4  
 gctctccctg ctccagcaag gacc atg agg gcg ctg gag ggg cca ggc ctg 51  
 Met Arg Ala Leu Glu Gly Pro Gly Leu  
 1 5

tgc ctg ctg tgc ctg gtg ttg gcg ctg cct gcc ctg ctg ccg gtg ccg 99  
 Ser Leu Leu Cys Leu Val Leu Ala Leu Pro Ala Leu Leu Pro Val Pro  
 10 15 20 25

gct gta cgc gga gtg gca gaa aca ccc acc tac ccc tgg cgg gac gca 147  
 Ala Val Arg Gly Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala  
 30 35 40

gag aca ggg gag cgg ctg gtg tgc gcc cag tgc ccc cca ggc acc ttt 195  
 Glu Thr Gly Glu Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe  
 45 50 55

gtg cag cgg ccg tgc cgc cga gac agc ccc acg acg tgt ggc ccg tgt 243  
 Val Gln Arg Pro Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys  
 60 65 70

cca ccg cgc cac tac acg cag ttc tgg aac tac ctg gag cgc tgc cgc 291  
 Pro Pro Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg  
 75 80 85

tac tgc aac gtc ctc tgc ggg gag cgt gag gag gag gca cgg gct tgc 339  
 Tyr Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys  
 90 95 100 105

cac gcc acc cac aac cgt gcc tgc cgc tgc cgc acc ggc ttc ttc gcg 387  
 His Ala Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala  
 110 115 120

cac gct ggt ttc tgc ttg gag cac gca tcg tgt cca cct ggt gcc ggc 435  
 His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly  
 125 130 135

gtg att gcc ccg ggc acc ccc agc cag aac acg cag tgc cag ccg tgc 483  
 Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys  
 140 145 150

```

ccc cca ggc acc ttc tca gcc agc agc tcc agc tca gag cag tgc cag 531
Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser Glu Gln Cys Gln
    155                160                165

ccc cac cgc aac tgc acg gcc ctg ggc ctg gcc ctc att gtg cca ggc 579
Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala Leu Ile Val Pro Gly
170                175                180                185

tct tcc tcc cat gac acc ctg tgc acc agc tgc act ggc ttc ccc ctc 627
Ser Ser Ser His Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu
                190                195                200

agc acc agg gta cca gga gct gag gag tgt gag cgt gcc gtc atc gac 675
Ser Thr Arg Val Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp
                205                210                215

ttt gtg gct ttc cag gac atc tcc atc aag agg ctg cag cgg ctg ctg 723
Phe Val Ala Phe Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu
    220                225                230

cag gcc ctc gag gcc ccg gag ggc tgg gct ccg aca cca agg gcg ggc 771
Gln Ala Leu Glu Ala Pro Glu Gly Trp Ala Pro Thr Pro Arg Ala Gly
    235                240                245

cgc gcg gcc ttg cag ctg aag ctg cgt cgg cgg ctc acg gag ctc ctg 819
Arg Ala Ala Leu Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu
250                255                260                265

ggg gcg cag gac ggg gcg ctg ctg gtg cgg ctg ctg cag gcg ctg cgc 867
Gly Ala Gln Asp Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg
    270                275                280

gtg gcc agg atg ccc ggg ctg gag cgg agc gtc cgt gag cgc ttc ctc 915
Val Ala Arg Met Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu
    285                290                295

cct gtg cac tgatcctggc cc 936
Pro Val His
    300

```

&lt;210&gt; 5

&lt;211&gt; 45

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 5

gagctagcca ccatgagggc gctggagggg ccaggcctgt cgctg

45

&lt;210&gt; 6

-7-

&lt;211&gt; 52

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 6

gtctcgttgt cccgccatgg gtaggtgggt gtttctgcc a ctccggtac ag 52

&lt;210&gt; 7

&lt;211&gt; 52

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;400&gt; 7

ggcagaaaca cccacctacc catggcgagg caacgagaca ggggagcggc tg 52

&lt;210&gt; 8

&lt;211&gt; 49

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 8

gtcgaatgacg gcacgctcac actcctcagc tcctgggtacc ctgggtgctg 49



## INTERNATIONAL SEARCH REPORT

International Application No.

PCT/US 02/00509

A. CLASSIFICATION OF SUBJECT MATTER  
IPC 7 C07K14/705 A61K38/17

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C07K C12N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, MEDLINE, BIOSIS, CHEM ABS Data, SEQUENCE SEARCH

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 00 58465 A (BECKER GERALD WAYNE ;COHEN FREDRIC JAY (US); GONZALEZ DEWHITT PATR) 5 October 2000 (2000-10-05) cited in the application page 31, paragraph 1 -page 32, paragraph 1; examples 8,14,15	1-20
Y	WO 00 58466 A (MICANOVIC RADMILA ;LILLY CO ELI (US); RATHNACHALAM RADHAKRISHNAN ( ) 5 October 2000 (2000-10-05) cited in the application page 13, paragraph 10; examples 1-9 page 24, paragraph 4 -page 26, paragraph 2 -/-	1-20

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

\* Special categories of cited documents:

\*A\* document defining the general state of the art which is not considered to be of particular relevance

\*E\* earlier document but published on or after the international filing date

\*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

\*O\* document referring to an oral disclosure, use, exhibition or other means

\*P\* document published prior to the international filing date but later than the priority date claimed

\*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

\*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

\*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

\*G\* document member of the same patent family

Date of the actual completion of the international search

10 September 2002

Date of mailing of the international search report

20/09/2002

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2  
NL - 2280 HV Rijswijk  
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,  
Fax: (+31-70) 340-3016

Authorized officer

Gurdjian, D

## INTERNATIONAL SEARCH REPORT

International Application No  
PCT/US 02/00509

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	KRONMAN C ET AL: "Involvement of oligomerization, N-glycosylation and sialylation in the clearance of cholinesterases from the circulation." THE BIOCHEMICAL JOURNAL. ENGLAND 1 NOV 1995, vol. 311 ( Pt 3), 1 November 1995 (1995-11-01), pages 959-967, XP008007650 ISSN: 0264-6021 abstract	1-20
A	PITTI ET AL: "Genomic amplification of a decoy receptor for FAS ligand in lung and colon cancer" NATURE, MACMILLAN JOURNALS LTD. LONDON, GB, vol. 396, 17 December 1998 (1998-12-17), pages 699-703, XP002139977 ISSN: 0028-0836 abstract; figure 1	1-20
A	YU K-Y ET AL: "A newly identified member of tumor necrosis factor receptor superfamily (TR6) suppresses LIGHT-mediated apoptosis" JOURNAL OF BIOLOGICAL CHEMISTRY, AMERICAN SOCIETY OF BIOLOGICAL CHEMISTS, BALTIMORE, MD, US, vol. 274, no. 20, 14 May 1999 (1999-05-14), pages 13733-13736, XP002161940 ISSN: 0021-9258 figure 1	1-20
A	BAI C ET AL: "OVEREXPRESSION OF M68/DCR3 IN HUMAN GASTROINTESTINAL TRACT TUMORS INDEPENDENT OF GENE AMPLIFICATION AND ITS LOCATION IN A FOUR-GENE CLUSTER" PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF USA, NATIONAL ACADEMY OF SCIENCE. WASHINGTON, US, vol. 97, no. 3, 1 February 2000 (2000-02-01), pages 1230-1235, XP002938755 ISSN: 0027-8424 abstract; figure 6	1-20
P,X	WO 01 18055 A (TIAN YU ;LILLY CO ELI (US); ATKINSON PAUL ROBERT (US); WITCHER DER) 15 March 2001 (2001-03-15) page 11, paragraph 2 -page 12, line 32	1-20

## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US 02/00509

### Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☒ Claims Nos.: 1-20  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:  
see FURTHER INFORMATION sheet PCT/ISA/210
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

### Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

#### Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

## FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Continuation of Box I.2

Claims Nos.: 1-20

Present claims 1-20 relate to a compound defined (inter alia) by reference to the following parameter : 'said composition having an average sialic acid content of about ... sialic acids per molecule of FLINT analog ' .

The use of these parameter in the present context is considered to lead to a lack of clarity within the meaning of Article 6 PCT. It is impossible to compare the parameter the applicant has chosen to employ with what is set out in the prior art. The lack of clarity is such as to render a meaningful complete search impossible. Consequently, the search has been restricted to FLINT analogs and the general concept of sialylation .

The applicant's attention is drawn to the fact that claims, or parts of claims, relating to inventions in respect of which no international search report has been established need not be the subject of an international preliminary examination (Rule 66.1(e) PCT). The applicant is advised that the EPO policy when acting as an International Preliminary Examining Authority is normally not to carry out a preliminary examination on matter which has not been searched. This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure.

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/US 02/00509

Patent document cited in search report		Publication date	Patent family member(s)	Publication date
WO 0058465	A	05-10-2000	AU 3739400 A	16-10-2000
			AU 3739500 A	16-10-2000
			EP 1165780 A2	02-01-2002
			EP 1165781 A2	02-01-2002
			WO 0058465 A2	05-10-2000
			WO 0058466 A2	05-10-2000
WO 0058466	A	05-10-2000	AU 3739400 A	16-10-2000
			AU 3739500 A	16-10-2000
			EP 1165780 A2	02-01-2002
			EP 1165781 A2	02-01-2002
			WO 0058465 A2	05-10-2000
			WO 0058466 A2	05-10-2000
WO 0118055	A	15-03-2001	AU 6891800 A	10-04-2001
			WO 0118055 A1	15-03-2001